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Ile Thr Thr Phe Met Gln Ser Glu Ile Val Ser Ser Asn Asp Ala Ile
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Thr Ser Tyr Asn Lys Asp Lys Ile Leu Asn Asp Ile Asn Ile Gln Ser
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Phe Gln Arg Leu Asn Ile Pro Phe Pro Cys Asp Cys Ile Gly Gly Glu
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 Asn Asn Phe Ser Leu Asp Asn Lys Ile Gly Gln Gly Gly Phe Gly Ala
 325 330 335
 Val Tyr Tyr Ala Glu Leu Arg Gly Lys Lys Thr Ala Ile Lys Lys Met
 340 345 350
 Asp Val Gln Ala Ser Thr Glu Phe Leu Cys Glu Leu Lys Val Leu Thr
 355 360 365

P200300646WO SEQ ID list

His Val His His Leu Asn Leu Val Arg Leu Ile Gly Tyr Cys Val Glu
 370 375 380
 Gly Ser Leu Phe Leu Val Tyr Glu His Ile Asp Asn Gly Asn Leu Gly
 385 390 395 400
 Gln Tyr Leu His Gly Ser Gly Lys Glu Pro Leu Pro Trp Ser Ser Arg
 405 410 415
 Val Gln Ile Ala Leu Asp Ala Ala Arg Gly Leu Glu Tyr Ile His Glu
 420 425 430
 His Thr Val Pro Val Tyr Ile His Arg Asp Val Lys Ser Ala Asn Ile
 435 440 445
 Leu Ile Asp Lys Asn Leu Arg Gly Lys Val Ala Asp Phe Gly Leu Thr
 450 455 460
 Lys Leu Ile Glu Val Gly Asn Ser Thr Leu Gln Thr Arg Leu Val Gly
 465 470 475 480
 Thr Phe Gly Tyr Met Pro Pro Glu Tyr Ala Gln Tyr Gly Asp Ile Ser
 485 490 495
 Pro Lys Ile Asp Val Tyr Ala Phe Gly Val Val Leu Phe Glu Leu Ile
 500 505 510
 Ser Ala Lys Asn Ala Val Leu Lys Thr Gly Glu Leu Val Ala Glu Ser
 515 520 525
 Lys Gly Leu Val Ala Leu Phe Glu Glu Ala Leu Asn Lys Ser Asp Pro
 530 535 540
 Cys Asp Ala Leu Arg Lys Leu Val Asp Pro Arg Leu Gly Glu Asn Tyr
 545 550 555 560
 Pro Ile Asp Ser Val Leu Lys Ile Ala Gln Leu Gly Arg Ala Cys Thr
 565 570 575
 Arg Asp Asn Pro Leu Leu Arg Pro Ser Met Arg Ser Leu Val Val Ala
 580 585 590
 Leu Met Thr Leu Ser Ser Leu Thr Glu Asp Cys Asp Asp Glu Ser Ser
 595 600 605
 Tyr Glu Ser Gln Thr Leu Ile Asn Leu Leu Ser Val Arg
 610 615 620

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 <211> 623

P200300646WO SEQ ID list

<212> PRT

<213> Lotus japonicus Gifu

<400> 25

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Ala Leu Ala Ser Tyr Tyr Ile Leu Pro Gly Val Phe Ile Leu Gln Asn
 35 40 45

Ile Thr Thr Phe Met Gln Ser Glu Ile Val Ser Ser Asn Asp Ala Ile
 50 55 60

Thr Ser Tyr Asn Lys Asp Lys Ile Leu Asn Asp Ile Asn Ile Gln Ser
 65 70 75 80

Phe Gln Arg Leu Asn Ile Pro Phe Pro Cys Asp Cys Ile Gly Gly Glu
 85 90 95

Phe Leu Gly His Val Phe Glu Tyr Ser Ala Ser Lys Gly Asp Thr Tyr
 100 105 110

Glu Thr Ile Ala Asn Leu Tyr Tyr Ala Asn Leu Thr Thr Val Asp Leu
 115 120 125

Leu Lys Arg Phe Asn Ser Tyr Asp Pro Lys Asn Ile Pro Val Asn Ala
 130 135 140

Lys Val Asn Val Thr Val Asn Cys Ser Cys Gly Asn Ser Gln Val Ser
 145 150 155 160

Lys Asp Tyr Gly Leu Phe Ile Thr Tyr Pro Ile Arg Pro Gly Asp Thr
 165 170 175

Leu Gln Asp Ile Ala Asn Gln Ser Ser Leu Asp Ala Gly Leu Ile Gln
 180 185 190

Ser Phe Asn Pro Ser Val Asn Phe Ser Lys Asp Ser Gly Ile Ala Phe
 195 200 205

Ile Pro Gly Arg Tyr Lys Asn Gly Val Tyr Val Pro Leu Tyr His Arg
 210 215 220

Thr Ala Gly Leu Ala Ser Gly Ala Ala Val Gly Ile Ser Ile Ala Gly
 225 230 235 240

Thr Phe Val Leu Leu Leu Ala Phe Cys Met Tyr Val Arg Tyr Gln
 245 250 255

P200300646WO SEQ ID list

Lys Lys Glu Glu Glu Lys Ala Lys Leu Pro Thr Asp Ile Ser Met Ala
 260 265 270
 Leu Ser Thr Gln Asp Ala Gly Asn Ser Ser Ser Ala Glu Tyr Glu Thr
 275 280 285
 Ser Gly Ser Ser Gly Pro Gly Thr Ala Ser Ala Thr Gly Leu Thr Ser
 290 295 300
 Ile Met Val Ala Lys Ser Met Glu Phe Ser Tyr Gln Glu Leu Ala Lys
 305 310 315 320
 Ala Thr Asn Asn Phe Ser Leu Asp Asn Lys Ile Gly Gln Gly Gly Phe
 325 330 335
 Gly Ala Val Tyr Tyr Ala Glu Leu Arg Gly Lys Lys Thr Ala Ile Lys
 340 345 350
 Lys Met Asp Val Gln Ala Ser Thr Glu Phe Leu Cys Glu Leu Lys Val
 355 360 365
 Leu Thr His Val His His Leu Asn Leu Val Arg Leu Ile Gly Tyr Cys
 370 375 380
 Val Glu Gly Ser Leu Phe Leu Val Tyr Glu His Ile Asp Asn Gly Asn
 385 390 395 400
 Leu Gly Gln Tyr Leu His Gly Ser Gly Lys Glu Pro Leu Pro Trp Ser
 405 410 415
 Ser Arg Val Gln Ile Ala Leu Asp Ala Ala Arg Gly Leu Glu Tyr Ile
 420 425 430
 His Glu His Thr Val Pro Val Tyr Ile His Arg Asp Val Lys Ser Ala
 435 440 445
 Asn Ile Leu Ile Asp Lys Asn Leu Arg Gly Lys Val Ala Asp Phe Gly
 450 455 460
 Leu Thr Lys Leu Ile Glu Val Gly Asn Ser Thr Leu Gln Thr Arg Leu
 465 470 475 480
 Val Gly Thr Phe Gly Tyr Met Pro Pro Glu Tyr Ala Gln Tyr Gly Asp
 485 490 495
 Ile Ser Pro Lys Ile Asp Val Tyr Ala Phe Gly Val Val Leu Phe Glu
 500 505 510
 Leu Ile Ser Ala Lys Asn Ala Val Leu Lys Thr Gly Glu Leu Val Ala
 515 520 525

P200300646WO SEQ ID list

Glu Ser Lys Gly Leu Val Ala Leu Phe Glu Glu Ala Leu Asn Lys Ser
 530 535 540

Asp Pro Cys Asp Ala Leu Arg Lys Leu Val Asp Pro Arg Leu Gly Glu
 545 550 555 560

Asn Tyr Pro Ile Asp Ser Val Leu Lys Ile Ala Gln Leu Gly Arg Ala
 565 570 575

Cys Thr Arg Asp Asn Pro Leu Leu Arg Pro Ser Met Arg Ser Leu Val
 580 585 590

Val Ala Leu Met Thr Leu Ser Ser Leu Thr Glu Asp Cys Asp Asp Glu
 595 600 605

Ser Ser Tyr Glu Ser Gln Thr Leu Ile Asn Leu Leu Ser Val Arg
 610 615 620

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 <213> Lotus japonicus

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<210> 27
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<210> 29
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 <212> DNA
 <213> Lotus japonicus

<400> 29
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<210> 30
 <211> 1853
 <212> DNA
 <213> Lotus filicaulis

<400> 30
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ttgtgaaaca	tatgtgacat	acacagctca	gtctccaaat	cttctgagcc	tgacaaacat	240
atctgatata	tttgatatca	gtcctttgtc	cattgcaaga	gccagtaaca	tagatgcagg	300
gaaggacaag	ctggttccag	gccaagtctt	actggtacct	gtaacttgcg	gttgcgccgg	360
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tgcaaccact	ttatatgaga	accttacaaa	ttggaatata	gtacaagctt	caaaccagg	480
ggtaaatacca	tatttggtgc	cagagcgcg	caaagtcgta	ttccctttat	tctgcagggtg	540
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caatgacaat	gtttcccttg	tgagtgccaa	gtttggtgca	tccccagcgg	acatattgac	660
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tcaaaggata	agcatagcag	tggatgttgc	tgtgggtctg	caatacatgc	atgaacatac	1320
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gacaaccaag	gagaacggcg	aggtggttat	gctgtggaag	gatatgtggg	agatctttga	1560
catagaagag	aatagagagg	agaggatcag	aaaatggatg	gacctaatt	tagagagctt	1620
ttatcatata	gataatgctc	tcagcttggc	atccttagca	gtgaattgca	cagctgataa	1680
gtctttgtct	cgaccctcca	tggctgaaat	tgttcttagc	ctctcctttc	tcactcaaca	1740
atcatctaac	cccacattag	agagatcctt	gacttcttct	gggttagatg	tagaagatga	1800
tgctcatatt	gtcacttcca	ttacagcacg	ttaagcaagg	gaaggtaatt	cag	1853

<210> 31
 <211> 595
 <212> PRT
 <213> Lotus filicaulis

<400> 31

P200300646WO SEQ ID list

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 Gly Pro Asp Phe Ser Cys Pro Val Asp Ser Pro Pro Ser Cys Glu Thr
 35 40 45
 Tyr Val Thr Tyr Thr Ala Gln Ser Pro Asn Leu Leu Ser Leu Thr Asn
 50 55 60
 Ile Ser Asp Ile Phe Asp Ile Ser Pro Leu Ser Ile Ala Arg Ala Ser
 65 70 75 80
 Asn Ile Asp Ala Gly Lys Asp Lys Leu Val Pro Gly Gln Val Leu Leu
 85 90 95
 Val Pro Val Thr Cys Gly Cys Ala Gly Asn His Ser Ser Ala Asn Thr
 100 105 110
 Ser Tyr Gln Ile Gln Lys Gly Asp Ser Tyr Asp Phe Val Ala Thr Thr
 115 120 125
 Leu Tyr Glu Asn Leu Thr Asn Trp Asn Ile Val Gln Ala Ser Asn Pro
 130 135 140
 Gly Val Asn Pro Tyr Leu Leu Pro Glu Arg Val Lys Val Val Phe Pro
 145 150 155 160
 Leu Phe Cys Arg Cys Pro Ser Lys Asn Gln Leu Asn Lys Gly Ile Gln
 165 170 175
 Tyr Leu Ile Thr Tyr Val Trp Lys Pro Asn Asp Asn Val Ser Leu Val
 180 185 190
 Ser Ala Lys Phe Gly Ala Ser Pro Ala Asp Ile Leu Thr Glu Asn Arg
 195 200 205
 Tyr Gly Gln Asp Phe Thr Ala Ala Thr Asn Leu Pro Ile Leu Ile Pro
 210 215 220
 Val Thr Gln Leu Pro Lys Leu Thr Gln Pro Ser Ser Asn Gly Arg Lys
 225 230 235 240
 Ser Ser Ile His Leu Leu Val Ile Leu Gly Ile Thr Leu Gly Cys Thr
 245 250 255
 Leu Leu Thr Ala Val Leu Thr Gly Thr Leu Val Tyr Val Tyr Cys Arg
 260 265 270

P200300646WO SEQ ID list

Arg Lys Lys Ala Leu Asn Arg Thr Ala Ser Ser Ala Glu Thr Ala Asp
 275 280 285
 Lys Leu Leu Ser Gly Val Ser Gly Tyr Val Ser Lys Pro Asn Val Tyr
 290 295 300
 Glu Ile Asp Glu Ile Met Glu Ala Thr Lys Asp Phe Ser Asp Glu Cys
 305 310 315 320
 Lys Val Gly Glu Ser Val Tyr Lys Ala Asn Ile Glu Gly Arg Val Val
 325 330 335
 Ala Val Lys Lys Ile Lys Glu Gly Gly Ala Asn Glu Glu Leu Lys Ile
 340 345 350
 Leu Gln Lys Val Asn His Gly Asn Leu Val Lys Leu Met Gly Val Ser
 355 360 365
 Ser Gly Tyr Asp Gly Asn Cys Phe Leu Val Tyr Glu Tyr Ala Glu Asn
 370 375 380
 Gly Ser Leu Ala Glu Trp Leu Phe Ser Lys Ser Ser Gly Thr Pro Asn
 385 390 395 400
 Ser Leu Thr Trp Ser Gln Arg Ile Ser Ile Ala Val Asp Val Ala Val
 405 410 415
 Gly Leu Gln Tyr Met His Glu His Thr Tyr Pro Arg Ile Ile His Arg
 420 425 430
 Asp Ile Thr Thr Ser Asn Ile Leu Leu Asp Ser Thr Phe Lys Ala Lys
 435 440 445
 Ile Ala Asn Phe Ala Met Ala Arg Thr Ser Thr Asn Pro Met Met Pro
 450 455 460
 Lys Ile Asp Val Phe Ala Phe Gly Val Leu Leu Ile Glu Leu Leu Thr
 465 470 475 480
 Gly Arg Lys Ala Met Thr Thr Lys Glu Asn Gly Glu Val Val Met Leu
 485 490 495
 Trp Lys Asp Met Trp Glu Ile Phe Asp Ile Glu Glu Asn Arg Glu Glu
 500 505 510
 Arg Ile Arg Lys Trp Met Asp Pro Asn Leu Glu Ser Phe Tyr His Ile
 515 520 525
 Asp Asn Ala Leu Ser Leu Ala Ser Leu Ala Val Asn Cys Thr Ala Asp
 530 535 540

P200300646WO SEQ ID list

Lys Ser Leu Ser Arg Pro Ser Met Ala Glu Ile Val Leu Ser Leu Ser
545 550 555 560

Phe Leu Thr Gln Gln Ser Ser Asn Pro Thr Leu Glu Arg Ser Leu Thr
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Ser Ser Gly Leu Asp Val Glu Asp Asp Ala His Ile Val Thr Ser Ile
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Thr Ala Arg
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<212> PRT
<213> Medicago truncatula

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<400> 32

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Ser Glu Thr Asn Phe Thr Cys Pro Val Asp Ser Pro Pro Ser Cys Glu
35 40 45

Thr Tyr Val Ala Tyr Arg Ala Gln Ser Pro Asn Phe Leu Ser Leu Ser
50 55 60

Asn Ile Ser Asp Ile Phe Asn Leu Ser Pro Leu Arg Ile Ala Lys Ala
65 70 75 80

Ser Asn Ile Glu Ala Glu Asp Lys Lys Leu Ile Pro Asp Gln Leu Leu
85 90 95

Leu Val Pro Val Thr Cys Gly Cys Thr Lys Asn His Ser Phe Ala Asn
100 105 110

Ile Thr Tyr Ser Ile Lys Gln Gly Asp Asn Phe Phe Ile Leu Ser Ile
115 120 125

Thr Ser Tyr Gln Asn Leu Thr Asn Tyr Leu Glu Phe Lys Asn Phe Asn
130 135 140

Pro Asn Leu Ser Pro Thr Leu Leu Pro Leu Asp Thr Lys Val Ser Val
145 150 155 160

P200300646WO SEQ ID list

Pro Leu Phe Cys Lys Cys Pro Ser Lys Asn Gln Leu Asn Lys Gly Ile
 165 170 175
 Lys Tyr Leu Ile Thr Tyr Val Trp Gln Asp Asn Asp Asn Val Thr Leu
 180 185 190
 Val Ser Ser Lys Phe Gly Ala Ser Gln Val Glu Met Leu Ala Glu Asn
 195 200 205
 Asn His Asn Phe Thr Ala Ser Thr Asn Arg Ser Val Leu Ile Pro Val
 210 215 220
 Thr Ser Leu Pro Lys Leu Asp Gln Pro Ser Ser Asn Gly Arg Lys Ser
 225 230 235 240
 Ser Ser Gln Asn Leu Ala Leu Ile Ile Gly Ile Ser Leu Gly Ser Ala
 245 250 255
 Phe Phe Ile Leu Val Leu Thr Leu Ser Leu Val Tyr Val Tyr Cys Leu
 260 265 270
 Lys Met Lys Arg Leu Asn Arg Ser Thr Ser Ser Ser Glu Thr Ala Asp
 275 280 285
 Lys Leu Leu Ser Gly Val Ser Gly Tyr Val Ser Lys Pro Thr Met Tyr
 290 295 300
 Glu Ile Asp Ala Ile Met Glu Gly Thr Thr Asn Leu Ser Asp Asn Cys
 305 310 315 320
 Lys Ile Gly Glu Ser Val Tyr Lys Ala Asn Ile Asp Gly Arg Val Leu
 325 330 335
 Ala Val Lys Lys Ile Lys Lys Asp Ala Ser Glu Glu Leu Lys Ile Leu
 340 345 350
 Gln Lys Val Asn His Gly Asn Leu Val Lys Leu Met Gly Val Ser Ser
 355 360 365
 Asp Asn Asp Gly Asn Cys Phe Leu Val Tyr Glu Tyr Ala Glu Asn Gly
 370 375 380
 Ser Leu Glu Glu Trp Leu Phe Ser Glu Ser Ser Lys Thr Ser Asn Ser
 385 390 395 400
 Val Val Ser Leu Thr Trp Ser Gln Arg Ile Thr Ile Ala Met Asp Val
 405 410 415
 Ala Ile Gly Leu Gln Tyr Met His Glu His Thr Tyr Pro Arg Ile Ile
 420 425 430

P200300646WO SEQ ID list

His Arg Asp Ile Thr Thr Ser Asn Ile Leu Leu Gly Ser Asn Phe Lys
 435 440 445

Ala Lys Ile Ala Asn Phe Gly Met Ala Arg Thr Ser Thr Asn Ser Met
 450 455 460

Met Pro Lys Ile Asp Val Phe Ala Phe Gly Val Val Leu Ile Glu Leu
 465 470 475 480

Leu Thr Gly Lys Lys Ala Met Thr Thr Lys Glu Asn Gly Glu Val Val
 485 490 495

Ile Leu Trp Lys Asp Phe Trp Lys Ile Phe Asp Leu Glu Gly Asn Arg
 500 505 510

Glu Glu Arg Leu Arg Lys Trp Met Asp Pro Lys Leu Glu Ser Phe Tyr
 515 520 525

Pro Ile Asp Asn Ala Leu Ser Leu Ala Ser Leu Ala Val Asn Cys Thr
 530 535 540

Ala Asp Lys Ser Leu Ser Arg Pro Thr Ile Ala Glu Ile Val Leu Cys
 545 550 555 560

Leu Ser Leu Leu Asn Gln Pro Ser Ser Glu Pro Met Leu Glu Arg Ser
 565 570 575

Leu Thr Ser Gly Leu Asp Ala Glu Ala Thr His Val Val Thr Ser Val
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Ile Ala Arg
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 <223> Primer to amplify fragment of NPR5

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24

<210> 34
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P200300646WO SEQ ID list

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 <223> To amplify a fragment of NPR5
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 <223> NPR5 5'RACE primer
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 <223> NPR5 gene PCR primer
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 ccagtgatag tagactgcct cataactttc atttgttcac aacttcacat ca atg gct 178
 Met Ala
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 gtc ttc ttt gtt tct ctt act ctt ggt gct cag att ctt tat gtg gta 226
 Val Phe Phe Val Ser Leu Thr Leu Gly Ala Gln Ile Leu Tyr Val Val
 5 10 15
 ctc atg ttt ttc act tgt att gaa gct caa tca caa cag acc aat gga 274
 Leu Met Phe Phe Thr Cys Ile Glu Ala Gln Ser Gln Gln Thr Asn Gly
 20 25 30
 aca aac ttt tca tgc cct tcc aat tca cct cct tca tgt gaa aca tat 322
 Thr Asn Phe Ser Cys Pro Ser Asn Ser Pro Pro Ser Cys Glu Thr Tyr
 35 40 45 50
 gtg aca tac ata tcc cag tcg cca aat ttt ttg agt ctg acc agc gta 370
 Val Thr Tyr Ile Ser Gln Ser Pro Asn Phe Leu Ser Leu Thr Ser Val
 55 60 65
 tct aat ata ttt gac acg agt cct ttg tca att gcc aga gcc agc aac 418
 Ser Asn Ile Phe Asp Thr Ser Pro Leu Ser Ile Ala Arg Ala Ser Asn
 70 75 80
 tta cag cat gag gaa gac aag ttg att cca ggc caa gtc tta ctg ata 466
 Leu Gln His Glu Glu Asp Lys Leu Ile Pro Gly Gln Val Leu Leu Ile
 85 90 95
 cca gta acc tgt ggt tgc act gga aac cgc tct ttc gcc aac atc tcc 514
 Pro Val Thr Cys Gly Cys Thr Gly Asn Arg Ser Phe Ala Asn Ile Ser
 100 105 110
 tat gag atc aac caa ggt gat agc ttc tac ttt gtt gcg acc act tta 562
 Tyr Glu Ile Asn Gln Gly Asp Ser Phe Tyr Phe Val Ala Thr Thr Leu
 115 120 125 130
 tac cag aat ctc aca aat tgg cat gca gtg atg gat tta aac cca ggt 610
 Tyr Gln Asn Leu Thr Asn Trp His Ala Val Met Asp Leu Asn Pro Gly
 135 140 145
 cta agt caa ttt act ttg cca ata ggc atc caa gtt gta att cct tta 658
 Leu Ser Gln Phe Thr Leu Pro Ile Gly Ile Gln Val Val Ile Pro Leu
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 ttc tgc aag tgt cct tca aag aac cag ctg gat aga ggg ata aag tac 706
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180 185 190															
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Asn Lys Leu Gly Ala Ser Pro Gln Asp Ile Leu Ser Glu Asn Asn Tyr															
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ggt caa aat ttc act gcc gca agc aac ctt cca gtt ttg atc cca gtt	850														
Gly Gln Asn Phe Thr Ala Ala Ser Asn Leu Pro Val Leu Ile Pro Val															
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Thr Leu Leu Pro Asp Leu Ile Gln Ser Pro Ser Asp Gly Arg Lys His															
230 235 240															
aga att ggt ctt cca gtt ata att ggt atc agc ctg gga tgc aca cta	946														
Arg Ile Gly Leu Pro Val Ile Ile Gly Ile Ser Leu Gly Cys Thr Leu															
245 250 255															
ctg gtt gtg gtt tca gca ata tta ctg gtg tgt gta tgt tgt ctg aaa	994														
Leu Val Val Val Ser Ala Ile Leu Leu Val Cys Val Cys Cys Leu Lys															
260 265 270															
atg aag agt ttg aat agg agt gct tca tca gct gaa act gca gat aaa	1042														
Met Lys Ser Leu Asn Arg Ser Ala Ser Ser Ala Glu Thr Ala Asp Lys															
275 280 285 290															
cta ctt tct gga gtt tca ggc tat gta agt aag cct aca atg tat gaa	1090														
Leu Leu Ser Gly Val Ser Ser Gly Tyr Val Ser Lys Pro Thr Met Tyr Glu															
295 300 305															
act ggt gca ata ttg gaa gct act atg aac ctc agt gag cag tgc aag	1138														
Thr Gly Ala Ile Leu Glu Ala Thr Met Asn Leu Ser Glu Gln Cys Lys															
310 315 320															
att ggg gaa tca gtg tac aag gct aac ata gag ggt aag gtt tta gca	1186														
Ile Gly Glu Ser Val Tyr Lys Ala Asn Ile Glu Gly Lys Val Leu Ala															
325 330 335															
gta aaa aga ttc aag gaa gat gtc acg gag gag ctg aaa att ctg cag	1234														
Val Lys Arg Phe Lys Glu Asp Val Thr Glu Glu Leu Lys Ile Leu Gln															
340 345 350															
aag gtg aat cat gga aat ctg gtg aaa cta atg ggt gtc tca tca gat	1282														
Lys Val Asn His Gly Asn Leu Val Lys Leu Met Gly Val Ser Ser Asp															
355 360 365 370															
aat gat gga aat tgt ttt gtg gtt tat gaa tat gct gaa aat ggg tct	1330														
Asn Asp Gly Asn Cys Phe Val Val Tyr Glu Tyr Ala Glu Asn Gly Ser															
375 380 385															
ctt gaa gag ttg ctt ttc gcc aag tct tgt tca gag aca tca aac tca	1378														
Leu Glu Glu Trp Leu Phe Ala Lys Ser Cys Ser Glu Thr Ser Asn Ser															
390 395 400															
agg acc tcc ctt aca tgg tgc cag agg ata agc ata gca gtg gat gtt	1426														
Arg Thr Ser Leu Thr Trp Cys Gln Arg Ile Ser Ile Ala Val Asp Val															
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tca atg ggt ctg cag tac atg cat gaa cat gct tat cca aga ata gtc	1474														
Ser Met Gly Leu Gln Tyr Met His Glu His Ala Tyr Pro Arg Ile Val															
420 425 430															
cac agg gac atc aca agc agt aat atc ctt ctt gac tcc aac ttt aag	1522														
His Arg Asp Ile Thr Ser Ser Asn Ile Leu Leu Asp Ser Asn Phe Lys															

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435	440	445	450	
gcc aag ata gca aat ttc tcc atg gcc aga act ttt acc aac ccc atg				1570
Ala Lys Ile Ala Asn Phe Ser Met Ala Arg Thr Phe Thr Asn Pro Met	455	460	465	
atg tca aaa ata gat gta ttt gct ttt ggg gtg gtt ctg ata gaa ttg				1618
Met Ser Lys Ile Asp Val Phe Ala Phe Gly Val Val Leu Ile Glu Leu	470	475	480	
ctt act ggc agg aaa gcc atg aca acc aaa gaa aat ggt gag gtg gtt				1666
Leu Thr Gly Arg Lys Ala Met Thr Thr Lys Glu Asn Gly Glu Val Val	485	490	495	
atg ctg tgg aag gac att tgg aag atc ttt gat caa gaa gag aat aga				1714
Met Leu Trp Lys Asp Ile Trp Lys Ile Phe Asp Gln Glu Glu Asn Arg	500	505	510	
gag gag agg ctc aga aaa tgg atg gat cct aag tta gat aat tat tat				1762
Glu Glu Arg Leu Arg Lys Trp Met Asp Pro Lys Leu Asp Asn Tyr Tyr	515	520	525	530
cct att gat tat gct ctc agc ttg gcc tcc ttg gca gtg aat tgc act				1810
Pro Ile Asp Tyr Ala Leu Ser Leu Ala Ser Leu Ala Val Asn Cys Thr	535	540	545	
gca gac aag tct ttg tcc aga cca acc ata gca gaa att gtc ctt agt				1858
Ala Asp Lys Ser Leu Ser Arg Pro Thr Ile Ala Glu Ile Val Leu Ser	550	555	560	
ctc tcc ctt ctc act caa cca tct ccc gcg aca ctg gag aga tcc ttg				1906
Leu Ser Leu Thr Gln Pro Ser Pro Ala Thr Leu Glu Arg Ser Leu	565	570	575	
act tct tct gga tta gat gta gaa gct act caa att gtc act tcc atc				1954
Thr Ser Ser Gly Leu Asp Val Glu Ala Thr Gln Ile Val Thr Ser Ile	580	585	590	
tca gct cgt tgattgagtg aagccaatct agttttctcac atccaagatg				2003
Ser Ala Arg	595			
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agttttcaac atttagtggt tccatccctg ttgttcttta tgtttgaggt agagttcgta				2123
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 <212> PRT
 <213> Phaseolus vulgaris

<400> 40

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Asn Gly Thr Asn Phe Ser Cys Pro Ser Asn Ser Pro Pro Ser Cys Glu
 35 40 45

P200300646WO SEQ ID list

Thr Tyr Val Thr Tyr Ile Ser Gln Ser Pro Asn Phe Leu Ser Leu Thr
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 Ser Val Ser Asn Ile Phe Asp Thr Ser Pro Leu Ser Ile Ala Arg Ala
 65 70 75 80
 Ser Asn Leu Gln His Glu Glu Asp Lys Leu Ile Pro Gly Gln Val Leu
 85 90 95
 Leu Ile Pro Val Thr Cys Gly Cys Thr Gly Asn Arg Ser Phe Ala Asn
 100 105 110
 Ile Ser Tyr Glu Ile Asn Gln Gly Asp Ser Phe Tyr Phe Val Ala Thr
 115 120 125
 Thr Leu Tyr Gln Asn Leu Thr Asn Trp His Ala Val Met Asp Leu Asn
 130 135 140
 Pro Gly Leu Ser Gln Phe Thr Leu Pro Ile Gly Ile Gln Val Val Ile
 145 150 155 160
 Pro Leu Phe Cys Lys Cys Pro Ser Lys Asn Gln Leu Asp Arg Gly Ile
 165 170 175
 Lys Tyr Leu Ile Thr His Val Trp Gln Pro Asn Asp Asn Val Ser Phe
 180 185 190
 Val Ser Asn Lys Leu Gly Ala Ser Pro Gln Asp Ile Leu Ser Glu Asn
 195 200 205
 Asn Tyr Gly Gln Asn Phe Thr Ala Ala Ser Asn Leu Pro Val Leu Ile
 210 215 220
 Pro Val Thr Leu Leu Pro Asp Leu Ile Gln Ser Pro Ser Asp Gly Arg
 225 230 235 240
 Lys His Arg Ile Gly Leu Pro Val Ile Ile Gly Ile Ser Leu Gly Cys
 245 250 255
 Thr Leu Leu Val Val Val Ser Ala Ile Leu Leu Val Cys Val Cys Cys
 260 265 270
 Leu Lys Met Lys Ser Leu Asn Arg Ser Ala Ser Ser Ala Glu Thr Ala
 275 280 285
 Asp Lys Leu Leu Ser Gly Val Ser Gly Tyr Val Ser Lys Pro Thr Met
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 Tyr Glu Thr Gly Ala Ile Leu Glu Ala Thr Met Asn Leu Ser Glu Gln
 305 310 315 320

P200300646WO SEQ ID list

Cys Lys Ile Gly Glu Ser Val Tyr Lys Ala Asn Ile Glu Gly Lys Val
 325 330 335
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 Leu Gln Lys Val Asn His Gly Asn Leu Val Lys Leu Met Gly Val Ser
 355 360 365
 Ser Asp Asn Asp Gly Asn Cys Phe Val Val Tyr Glu Tyr Ala Glu Asn
 370 375 380
 Gly Ser Leu Glu Glu Trp Leu Phe Ala Lys Ser Cys Ser Glu Thr Ser
 385 390 395 400
 Asn Ser Arg Thr Ser Leu Thr Trp Cys Gln Arg Ile Ser Ile Ala Val
 405 410 415
 Asp Val Ser Met Gly Leu Gln Tyr Met His Glu His Ala Tyr Pro Arg
 420 425 430
 Ile Val His Arg Asp Ile Thr Ser Ser Asn Ile Leu Leu Asp Ser Asn
 435 440 445
 Phe Lys Ala Lys Ile Ala Asn Phe Ser Met Ala Arg Thr Phe Thr Asn
 450 455 460
 Pro Met Met Ser Lys Ile Asp Val Phe Ala Phe Gly Val Val Leu Ile
 465 470 475 480
 Glu Leu Leu Thr Gly Arg Lys Ala Met Thr Thr Lys Glu Asn Gly Glu
 485 490 495
 Val Val Met Leu Trp Lys Asp Ile Trp Lys Ile Phe Asp Gln Glu Glu
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 Asn Arg Glu Glu Arg Leu Arg Lys Trp Met Asp Pro Lys Leu Asp Asn
 515 520 525
 Tyr Tyr Pro Ile Asp Tyr Ala Leu Ser Leu Ala Ser Leu Ala Val Asn
 530 535 540
 Cys Thr Ala Asp Lys Ser Leu Ser Arg Pro Thr Ile Ala Glu Ile Val
 545 550 555 560
 Leu Ser Leu Ser Leu Leu Thr Gln Pro Ser Pro Ala Thr Leu Glu Arg
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<210> 42
<211> 30
<212> DNA
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<220>
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<223> Primer to amplify NPR5 gene fragment

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<210> 43
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<212> DNA
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<220>
<221> misc_feature
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<223> NPR5 5'RACE primer

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<210> 44
<211> 19
<212> DNA
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<223> NPR5 3'RACE primer

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P200300646WO SEQ ID list

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 Met Ala Val Phe Phe Pro Phe Leu Pro Leu His Ser Gln Ile
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 ctt tgt ctt gtg atc atg ttg ttt tcc act aat att gta gct caa tca 158
 Leu Cys Leu Val Ile Met Leu Phe Ser Thr Asn Ile Val Ala Gln Ser
 15 20 25 30
 caa cag gac aat aga aca aac ttt tca tgc cct tct gat tca ccg cct 206
 Gln Gln Asp Asn Arg Thr Asn Phe Ser Cys Pro Ser Asp Ser Pro Pro
 35 40 45
 tca tgt gaa acc tat gta aca tac att gct cag tct cca aat ttt ttg 254
 Ser Cys Glu Thr Tyr Val Thr Tyr Ile Ala Gln Ser Pro Asn Phe Leu
 50 55 60
 agt cta acc aac ata tcc aat ata ttt gac aca agc cct tta tcc att 302
 Ser Leu Thr Asn Ile Ser Asn Ile Phe Asp Thr Ser Pro Leu Ser Ile
 65 70 75
 gca aga gcc agt aac tta gag cct atg gat gac aag cta gtc aaa gac 350

P200300646WO SEQ ID list

Ala Arg Ala Ser Asn Leu Glu Pro Met Asp Asp Lys Leu Val Lys Asp	
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caa gtc tta ctc gta cca gta acc tgt ggt tgc act gga aac cgc tct	398
Gln Val Leu Leu Val Pro Val Thr Cys Gly Cys Thr Gly Asn Arg Ser	
95 100 105 110	
ttt gcc aat atc tcc tat gag atc aac caa ggt gat agc ttc tac ttt	446
Phe Ala Asn Ile Ser Tyr Glu Ile Asn Gln Gly Asp Ser Phe Tyr Phe	
115 120 125	
gtt gca acc act tca tac gag aat ctc acg aat tgg cgt gca gtg atg	494
Val Ala Thr Thr Ser Tyr Glu Asn Leu Thr Asn Trp Arg Ala Val Met	
130 135 140	
gat tta aac ccc gtt cta agt cca aat aag ttg cca ata gga atc caa	542
Asp Leu Asn Pro Val Leu Ser Pro Asn Lys Leu Pro Ile Gly Ile Gln	
145 150 155	
gta gta ttt cct tta ttc tgc aag tgc cct tca aag aac cag ttg gac	590
Val Val Phe Pro Leu Phe Cys Lys Cys Pro Ser Lys Asn Gln Leu Asp	
160 165 170	
aaa gag ata aag tac ctg att aca tac gtg tgg aag ccc ggt gac aat	638
Lys Glu Ile Lys Tyr Leu Ile Thr Tyr Val Trp Lys Pro Gly Asp Asn	
175 180 185 190	
gtt tcc ctt gta agt gac aag ttt ggt gca tca cca gag gac ata atg	686
Val Ser Leu Val Ser Asp Lys Phe Gly Ala Ser Pro Glu Asp Ile Met	
195 200 205	
agt gaa aac aac tat ggt cag aac ttt act gct gca aac aac ctt cca	734
Ser Glu Asn Asn Tyr Gly Gln Asn Phe Thr Ala Ala Asn Asn Leu Pro	
210 215 220	
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Val Leu Ile Pro Val Thr Arg Leu Pro Val Leu Ala Arg Ser Pro Ser	
225 230 235	
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Asp Gly Arg Lys Gly Gly Ile Arg Leu Pro Val Ile Ile Gly Ile Ser	
240 245 250	
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Leu Gly Cys Thr Leu Leu Val Leu Val Leu Ala Val Leu Leu Val Tyr	
255 260 265 270	
gta tat tgt ctg aaa atg aag act ttg aat agg agt gct tca tcg gct	926
Val Tyr Cys Leu Lys Met Lys Thr Leu Asn Arg Ser Ala Ser Ser Ala	
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Glu Thr Ala Asp Lys Leu Leu Ser Gly Val Ser Gly Tyr Val Ser Lys	
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Pro Thr Met Tyr Glu Thr Asp Ala Ile Met Glu Ala Thr Met Asn Leu	
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agt gag cag tgc aag att ggg gaa tca gtg tac aag gca aac ata gag	1070
Ser Glu Gln Cys Lys Ile Gly Glu Ser Val Tyr Lys Ala Asn Ile Glu	
320 325 330	
ggg aag gtt ttg gca gta aaa aga ttc aag gaa gat gtc acg gaa gag	1118
Gly Lys Val Leu Ala Val Lys Arg Phe Lys Glu Asp Val Thr Glu Glu	
335 340 345 350	
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P200300646WO SEQ ID list

Leu Lys Ile Leu	Gln 355	Lys Val	Asn His	Gly 360	Asn Leu Val	Lys	Leu Met 365	
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Gly Val Ser Ser	370	Asp Asn Asp	Gly Asn 375	Cys Phe Val	Val	Tyr 380	Glu Tyr	
gct gaa aat ggg tct ctt gat gag tgg cta ttc tcc aag tct tgt tca								1262
Ala Glu Asn 385	Gly Ser Leu Asp 390	Trp Leu Phe Ser 395	Lys Ser Cys Ser					
gac aca tca aac tca agg gca tcc ctt aca tgg tgt cag agg ata agc								1310
Asp Thr 400	Asn Ser Arg Ala Ser Leu Thr 405	Trp Cys 410	Gln Arg Ile Ser					
atg gca gtg gat gtt gcg atg ggt ttg cag tac atg cat gaa cat gct								1358
Met Ala Val Asp Val 415	Ala Met Gly Leu Gln 420	Tyr Met His Glu His 430						
tat cca aga ata gtc cac agg gac atc aca agc agt aat atc ctt ctt								1406
Tyr Pro Arg Ile 435	His Arg Asp Ile 440	Thr Ser Ser Asn Ile 445	Leu Leu					
gac tcg aac ttt aag gcc aag ata gca aat ttc tcc atg gcc aga act								1454
Asp Ser Asn Phe Lys Ala Lys Ile 450	Ala Asn Phe Ser Met 455	Ala Arg Thr 460						
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Phe Thr Asn Pro Met Met Pro Lys Ile Asp Val Phe Ala Phe Gly Val 465								
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Val Leu Ile Glu Leu Leu Thr Gly Arg Lys Ala Met Thr Thr Lys Glu 480								
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Asn Gly Glu Val Val Met Leu Trp Lys Asp Ile Trp Lys Ile Phe Asp 495								
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Gln Glu Glu Asn Arg Glu Glu Arg Leu Lys Lys Trp Met Asp Pro Lys 515								
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Leu Glu Ser Tyr 530	Pro Ile Asp Tyr 535	Ala Leu Ser Leu 540						
gcg gtg aat tgt act gca gat aag tct ttg tcc aga cca acc att gca								1742
Ala Val Asn Cys Thr Ala Asp Lys Ser Leu Ser Arg Pro Thr Ile Ala 545								
gaa att gtc ctt agc ctc tcc ctt ctc act caa cca tct ccc gca aca								1790
Glu Ile Val Leu Ser Leu Ser Leu Leu Thr Gln Pro Ser Pro Ala Thr 560								
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Leu Glu Arg Ser Leu Thr Ser Ser Gly Leu Asp Val Glu Ala Thr Gln 575								
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Ile Val Thr Ser Ile Ala Ala Arg 595								
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 <212> PRT
 <213> Glycine max

<400> 48

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Asp Asn Arg Thr Asn Phe Ser Cys Pro Ser Asp Ser Pro Pro Ser Cys
 35 40 45

Glu Thr Tyr Val Thr Tyr Ile Ala Gln Ser Pro Asn Phe Leu Ser Leu
 50 55 60

Thr Asn Ile Ser Asn Ile Phe Asp Thr Ser Pro Leu Ser Ile Ala Arg
 65 70 75 80

Ala Ser Asn Leu Glu Pro Met Asp Asp Lys Leu Val Lys Asp Gln Val
 85 90 95

Leu Leu Val Pro Val Thr Cys Gly Cys Thr Gly Asn Arg Ser Phe Ala
 100 105 110

Asn Ile Ser Tyr Glu Ile Asn Gln Gly Asp Ser Phe Tyr Phe Val Ala
 115 120 125

Thr Thr Ser Tyr Glu Asn Leu Thr Asn Trp Arg Ala Val Met Asp Leu
 130 135 140

Asn Pro Val Leu Ser Pro Asn Lys Leu Pro Ile Gly Ile Gln Val Val
 145 150 155 160

Phe Pro Leu Phe Cys Lys Cys Pro Ser Lys Asn Gln Leu Asp Lys Glu
 165 170 175

Ile Lys Tyr Leu Ile Thr Tyr Val Trp Lys Pro Gly Asp Asn Val Ser
 180 185 190

Leu Val Ser Asp Lys Phe Gly Ala Ser Pro Glu Asp Ile Met Ser Glu
 195 200 205

Asn Asn Tyr Gly Gln Asn Phe Thr Ala Ala Asn Asn Leu Pro Val Leu
 210 215 220

Ile Pro Val Thr Arg Leu Pro Val Leu Ala Arg Ser Pro Ser Asp Gly
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P200300646WO SEQ ID list

500

505

510

Glu Asn Arg Glu Glu Arg Leu Lys Lys Trp Met Asp Pro Lys Leu Glu
515 520 525

Ser Tyr Tyr Pro Ile Asp Tyr Ala Leu Ser Leu Ala Ser Leu Ala Val
530 535 540

Asn Cys Thr Ala Asp Lys Ser Leu Ser Arg Pro Thr Ile Ala Glu Ile
545 550 555 560

Val Leu Ser Leu Ser Leu Leu Thr Gln Pro Ser Pro Ala Thr Leu Glu
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Thr Ser Ile Ala Ala Arg
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<212> DNA
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 1 5 10 15
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 Cys Ala Phe Phe Lys Val Asp Ser Lys Cys Val Lys Gly Cys Asp Leu
 20 25 30
 gct tta gct tct tac tat gta atg cct tta gtt gaa ctc cca act ata 206
 Ala Leu Ala Ser Tyr Tyr Val Met Pro Leu Val Glu Leu Pro Thr Ile
 35 40 45
 aaa aac tat atg caa tca aag ata gtt acc aac tct tct gat gtt tta 254
 Lys Asn Tyr Met Gln Ser Lys Ile Val Thr Asn Ser Ser Asp Val Leu
 50 55 60
 aat agt tac aac aaa gtc tta gta acc aat cat ggt aat att ttt tcc 302
 Asn Ser Tyr Asn Lys Val Leu Val Thr Asn His Gly Asn Ile Phe Ser
 65 70 75
 tat ttt aga atc aac att cca ttc cca tgt gaa tgt att gga ggt gag 350
 Tyr Phe Arg Ile Asn Ile Pro Phe Pro Cys Glu Cys Ile Gly Gly Glu
 80 85 90 95
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 Phe Leu Gly His Val Phe Glu Tyr Thr Thr Lys Lys Gly Asp Thr Tyr
 100 105 110
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 115 120 125
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 130 135 140
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 145 150 155
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 Lys Asp Tyr Gly Leu Phe Val Thr Tyr Pro Leu Arg Ser Thr Asp Ser
 160 165 170 175
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aaa Lys	aag Lys	gaa Glu	gaa Glu	gag Glu 260	aaa Lys	act Thr	ata Ile	ctg Leu	ccc Pro 265	caa Gln	gtt Val	tct Ser	aag Lys	gcg Ala 270	ctt Leu	878
tcg Ser	act Thr	caa Gln	gat Asp 275	ggg Gly	aat Asn	gcc Ala	tcg Ser	agt Ser 280	agt Ser	gga Gly	gaa Glu	tat Tyr	gaa Glu 285	act Thr	tca Ser	926
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gag Glu	ggg Gly 385	tcg Ser	ctt Leu	ttc Phe	ctt Leu	gtc Val 390	tat Tyr	gaa Glu	cat His	att Ile	gac Asp 395	aat Asn	gga Gly	aac Asn	ttg Leu	1262
ggg Gly 400	caa Gln	tat Tyr	ttg Leu	cat His	ggg Gly 405	aaa Lys	gat Asp	aaa Lys	gag Glu	cca Pro 410	tta Leu	cca Pro	tgg Trp	tct Ser	agt Ser 415	1310
aga Arg	gtc Val	caa Gln	att Ile	gct Ala 420	cta Leu	gat Asp	tca Ser	gca Ala	cga Arg 425	ggc Gly	ctt Leu	gaa Glu	tac Tyr	att Ile 430	cat His	1358
gaa Glu	cat His	acc Thr	gtg Val 435	cct Pro	gtg Val	tat Tyr	atc Ile	cat His 440	cgc Arg	gat Asp	gta Val	aaa Lys	tca Ser	gca Ala	aac Asn	1406
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acc Thr	aaa Lys 465	ctt Leu	att Ile	gaa Glu	gtt Val	gga Gly 470	aat Asn	tcc Ser	aca Thr	ctt Leu	cac His 475	act Thr	cgt Arg	ctt Leu	gtt Val	1502
gga Gly 480	act Thr	ttt Phe	gga Gly	tac Tyr	atg Met 485	cca Pro	cca Pro	gaa Glu	tat Tyr	gct Ala 490	caa Gln	tat Tyr	ggg Gly	gac Asp	gtt Val 495	1550
tct Ser	ccg Pro	aaa Lys	ata Ile	gac Asp	gta Val	tat Tyr	gct Ala	ttt Phe	gga Gly	gtt Val	gtt Val	ctt Leu	tat Tyr	gaa Glu	ctg Leu	1598

P200300646WO SEQ ID list

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aac tat cca att gat tct gtt tta aag atg gct caa ctt ggg aga gca Asn Tyr Pro Ile Asp Ser 565 Val Leu Lys Met Ala Gln Leu Gly Arg Ala 575			1790
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Asn Tyr Met Gln Ser Lys Ile Val Thr Asn Ser Ser Asp Val Leu Asn 50 55 60			
Ser Tyr Asn Lys Val Leu Val Thr Asn His Gly Asn Ile Phe Ser Tyr 65 70 75 80			
Phe Arg Ile Asn Ile Pro Phe Pro Cys Glu Cys Ile Gly Gly Glu Phe 85 90 95			

P200300646WO SEQ ID list

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 Lys Lys Phe Asn Ser Tyr Asp Pro Asn His Ile Pro Ala Lys Ala Lys
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 Val Asn Val Thr Val Asn Cys Ser Cys Gly Asn Ser Gln Ile Ser Lys
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 Asp Tyr Gly Leu Phe Val Thr Tyr Pro Leu Arg Ser Thr Asp Ser Leu
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 Glu Lys Ile Ala Asn Glu Ser Lys Leu Asp Glu Gly Leu Ile Gln Asn
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 Phe Asn Pro Asp Val Asn Phe Ser Arg Gly Ser Gly Ile Val Phe Ile
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 210 215 220
 Gly Val Gly Lys Gly Val Ala Ile Gly Ile Ser Ile Ala Gly Val Phe
 225 230 235 240
 Ala Val Leu Leu Phe Val Ile Cys Ile Tyr Val Lys Tyr Phe Gln Lys
 245 250 255
 Lys Glu Glu Glu Lys Thr Ile Leu Pro Gln Val Ser Lys Ala Leu Ser
 260 265 270
 Thr Gln Asp Gly Asn Ala Ser Ser Ser Gly Glu Tyr Glu Thr Ser Gly
 275 280 285
 Ser Ser Gly His Gly Thr Gly Ser Ala Ala Gly Leu Thr Gly Ile Met
 290 295 300
 Val Ala Lys Ser Thr Glu Phe Ser Tyr Gln Glu Leu Ala Lys Ala Thr
 305 310 315 320
 Asp Asn Phe Ser Leu Asp Asn Lys Ile Gly Gln Gly Gly Phe Gly Ala
 325 330 335
 Val Tyr Tyr Ala Glu Leu Arg Gly Glu Lys Thr Ala Ile Lys Lys Met
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 Asn Val Gln Ala Ser Ser Glu Phe Leu Cys Glu Leu Lys Val Leu Thr
 355 360 365

P200300646WO SEQ ID list

His Val His His Leu Asn Leu Val Arg Leu Ile Gly Tyr Cys Val Glu
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 Gly Ser Leu Phe Leu Val Tyr Glu His Ile Asp Asn Gly Asn Leu Gly
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 Gln Tyr Leu His Gly Lys Asp Lys Glu Pro Leu Pro Trp Ser Ser Arg
 405 410 415
 Val Gln Ile Ala Leu Asp Ser Ala Arg Gly Leu Glu Tyr Ile His Glu
 420 425 430
 His Thr Val Pro Val Tyr Ile His Arg Asp Val Lys Ser Ala Asn Ile
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 Leu Ile Asp Lys Asn Leu Arg Gly Lys Val Ala Asp Phe Gly Leu Thr
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 Thr Phe Gly Tyr Met Pro Pro Glu Tyr Ala Gln Tyr Gly Asp Val Ser
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 Pro Lys Ile Asp Val Tyr Ala Phe Gly Val Val Leu Tyr Glu Leu Ile
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 Ser Lys Gly Leu Val Ala Leu Phe Glu Lys Ala Leu Asn Gln Ile Asp
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 Pro Ser Glu Ala Leu Arg Lys Leu Val Asp Pro Arg Leu Lys Glu Asn
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 Tyr Pro Ile Asp Ser Val Leu Lys Met Ala Gln Leu Gly Arg Ala Cys
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 Thr Arg Asp Asn Pro Leu Leu Arg Pro Ser Met Arg Ser Leu Val Val
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Pro	Val	Lys	Ala	Lys	Val	Asn	Val	Thr	Val	Asn	Cys	Ser	Cys	Gly	Asn	
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Ser	Gln															

P200300646WO SEQ ID list

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P200300646WO SEQ ID list

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 Leu Val Gly Thr Phe Gly Tyr Met Pro Pro Glu Tyr Ala Gln Tyr Gly
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 Asp Val Ser Pro Lys Ile Asp Val Tyr Ala Phe Gly Val Val Leu Tyr
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 Leu Gly His Val Phe Glu Tyr Thr Ala Asn Glu Gly Asp Thr Tyr Asp
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 Leu Ile Ala Asn Thr Tyr Tyr Ala Ser Leu Thr Thr Val Glu Val Leu
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 370 375 380
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 Ile Asp Val Tyr Ala Phe Gly Val Val Leu Tyr Glu Leu Ile Ser Ala
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 Ala Leu Arg Lys Leu Val Asp Pro Arg Leu Lys Glu Asn Tyr Pro Ile
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Ser Leu Thr Asn Leu Leu Ser Val Arg
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